
Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2011; month=5; day=5; hr=10; min=21; sec=28; ms=567;]

Validated By CRFValidator v 1.0.3

Application No: 10598736 Version No: 2.0

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Finished: 2011-04-29 12:39:40.289

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Actual SeqID Count: 12

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<110> Lotersztajn, Sophie
 Mallat, Ariane
 Grenard, Pascale
 Julien, Boris
 Nhieu, Jeanne V.

<120> Use of Antagonists of the CBI Receptor for the Manufacture of a Composition Useful for the Treatment of Hepatic Diseases

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Pro Leu Thr Ser Phe Arg Gly Ser Pro Phe Gln Glu Lys Met Thr Ala 50 55 60

Gly Asp Asn Pro Gln Leu Val Pro Ala Asp Gln Val Asn Ile Thr Glu 65 70 75 80

Phe Tyr Asn Lys Ser Leu Ser Ser Phe Lys Glu Asn Glu Glu Asn Ile 85 90 95

Gln Cys Gly Glu Asn Phe Met Asp Ile Glu Cys Phe Met Val Leu Asn 100 105 110

Pro Ser Gln Gln Leu Ala Ile Ala Val Leu Ser Leu Thr Leu Gly Thr

115 120 125

Phe Thr Val Leu Glu Asn Leu Leu Val Leu Cys Val Ile Leu His Ser 130 135 140

Arg Ser Leu Arg Cys Arg Pro Ser Tyr His Phe Ile Gly Ser Leu Ala 145 150 150

Val Ala Asp Leu Leu Gly Ser Val Ile Phe Val Tyr Ser Phe Ile Asp

165
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Leu Gly Gly Val Thr Ala Ser Phe Thr Ala Ser Val Gly Ser Leu Phe
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Leu Thr Ala Ile Asp Arg Tyr Ile Ser Ile His Arg Pro Leu Ala Tyr 210 220

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Trp Thr Ile Ala Ile Val Ile Ala Val Leu Pro Leu Leu Gly Trp Asn
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Cys Glu Lys Leu Gln Ser Val Cys Ser Asp Ile Phe Pro His Ile Asp 260 270

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Thr Ser Glu Asp Gly Lys Val Gln Val Thr Arg Pro Asp Gln Ala Arg
325 330 335

Met Asp Ile Arg Leu Ala Lys Thr Leu Val Leu Ile Leu Val Val Leu 340 350

Ile Ile Cys Trp Gly Pro Leu Leu Ala Ile Met Val Tyr Asp Val Phe Gly Lys Met Asn Lys Leu Ile Lys Thr Val Phe Ala Phe Cys Ser Met Leu Cys Leu Leu Asn Ser Thr Val Asn Pro Ile Ile Tyr Ala Leu Arg Ser Lys Asp Leu Arg His Ala Phe Arg Ser Met Phe Pro Ser Cys Glu Gly Thr Ala Gln Pro Leu Asp Asn Ser Met Gly Asp Ser Asp Cys Leu His Lys His Ala Asn Asn Ala Ala Ser Val His Arg Ala Ala Glu Ser Cys Ile Lys Ser Thr Val Lys Ile Ala Lys Val Thr Met Ser Val Ser Thr Asp Thr Ser Ala Glu Ala Leu <210> 2 <211> <212> PRT <213> Homo sapiens <400> 2 Met Ala Leu Gln Ile Pro Pro Ser Ala Pro Ser Pro Leu Thr Ser Cys Thr Trp Ala Gln Met Thr Phe Ser Thr Lys Thr Ser Lys Glu Asn Glu Glu Asn Ile Gln Cys Gly Glu Asn Phe Met Asp Ile Glu Cys Phe Met

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Leu	His	Ser	Arg	Ser 85	Leu	Arg	Cys	Arg	Pro 90	Ser	Tyr	His	Phe	Ile 95	Gly
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Phe	Ile	Asp 115	Phe	His	Val	Phe	His 120	Arg	Lys	Asp	Ser	Arg 125	Asn	Val	Phe
Leu		_	Leu	_	_						Thr 140		Ser	Val	Gly
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Leu	Ala	Tyr	Lys	Arg 165	Ile	Val	Thr	Arg	Pro 170	Lys	Ala	Val	Val	Ala 175	Phe
Cys	Leu	Met	Trp 180	Thr	Ile	Ala	Ile	Val 185	Ile	Ala	Val	Leu	Pro 190	Leu	Leu
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Leu 225	Leu	Leu	Phe	Ile	Val 230	Tyr	Ala	Tyr	Met	Tyr 235	Ile	Leu	Trp	Lys	Ala 240
His	Ser	His	Ala	Val 245	Arg	Met	Ile	Gln	Arg 250	Gly	Thr	Gln	Lys	Ser 255	Ile
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290 295 300

Asp Val Phe Gly Lys Met Asn Lys Leu Ile Lys Thr Val Phe Ala Phe 305 310 315 320

Ala Leu Arg Ser Lys Asp Leu Arg His Ala Phe Arg Ser Met Phe Pro 340 345 350

Ser Cys Glu Gly Thr Ala Gln Pro Leu Asp Asn Ser Met Gly Asp Ser 355 360 365

Asp Cys Leu His Lys His Ala Asn Asn Ala Ala Ser Val His Arg Ala 370 380

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